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Twin-Arginine Translocation in Bacillus Bron et al. SN# 09/954,737 Docket No. GC634-2 Sheet 1 of 11

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TatA(Eco)	M-GGISIWQLLIIAVIVVLLFGTKKLG	- 26
TatE(Eco)	M-GEISITKLLVVAALVVLLFGTKKLR	- 26
TatAy(Bsu)	MPIGPGSLAVIAIVALIIFGPKKLP	- 25
TatAd(Bsu)	MFSNIGIPGLILIFVIAIIIFGPSKLP	
TatAc(Bsu)	M-ELSFTKILVILFVGFLVFGPDKLP	
TatB (Eco)	MF-DIGFSELLLVFIIGLVVLGPQRLPVAVKTVAGWIRALRSLATTVQNELTQELKLQ	
, ,	*	
m-+3 (D)		64
TatA(Eco)	PKQDKTSQDADFTAKTITLGGDLGAAIKGFKKAMNDDDA-AAKKGADVDLQAEKL	63
TatE(Eco)	ELGKAAGDTLREFKNATKGLTSDEEEKKKEDQ	57
TatAy(Bsu) TatAd(Bsu)	SDEEERREDQ	64
TatAc(Bsu)		57
TatB (Eco)	EFQDSLKKVEKASLTNLTPELKASMDELRQAAESMKRSYVANDPEKASDEAHTIHNP	114
Tach (ECO)	EFQDSDKKVEKASDINDIFEDKASNDEDKQARESNKKSIVANDFEKASDEANIIINF	114
TatA(Eco)	ADKQADTNQEQAKTEDAKRHDKEQV SHKE	89
TatE(Eco)	SHKE	67
TatAy(Bsu)		57
TatAd(Bsu)	QDKNAG	70
TatAc(Bsu)	EDKQM-	62
TatB (Eco)	VVKDNEAAHEGVTPAAAQTQASSPEQKPETTPEPVVKPAADAEPKTAAPSPSSSDKP	171
	FIG1A	
TatC (Eco)	MSVEDTOPI.TTHI.TELEKRI.INCTTAVIVIELCIVVEANDIVH-IVSAPI.IK	51
TatC (Eco)	MSVEDTQPLITHLIELRKRLLNCIIAVIVIFLCLVYFANDIYH-LVSAPLIK MTRMKVNOMSLLEHIAELRKRLLTVALAEVVEFTAGEFLAKETTVYLOETDEAK	51 50
TatCy (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK	50
-		
TatCy (Bsu) TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * *	50 51
TatCy (Bsu) TatCd (Bsu) TatC (Eco)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * ** * * * * * * QLPQGSTMIATDVASPFFTPIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR	50 51 105
TatCy (Bsu) TatCd (Bsu) TatC (Eco) TatCy (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * . * * * * * QLPQGSTMIATDVASPFFTPIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR QLTLNAFNLTDPLYVFMQFAFIIGIVLTSPVILYQLWAFVSPGLYEKERK	50 51 105 104
TatCy (Bsu) TatCd (Bsu) TatC (Eco)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * ** * * * * * * QLPQGSTMIATDVASPFFTPIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR	50 51 105
TatCy (Bsu) TatCd (Bsu) TatC (Eco) TatCy (Bsu) TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * * * * * * * * * * *	50 51 105 104 98
TatCy (Bsu) TatCd (Bsu) TatC (Eco) TatCy (Bsu) TatCd (Bsu) TatCd (Eco)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * *  QLPQGSTMIATDVASPFFTPIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR QLTLNAFNLTDPLYVFMQFAFIIGIVLTSPVILYQLWAFVSPGLYEKERKLAVLGPSEILWVYMMLSGICAIAASIPVAAYQLWRFVAPALTKTERK	50 51 105 104
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * . * * * *  QLPQGSTMIATDVASPFFTEIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR QLTLNAFNLTDPLYVFMQFAFIIGIVLTSPVILYQLWAFVSPGLYEKERKLAVLGPSEILWVYMMLSGICAIAASIPVAAYQLWRFVAPALTKTERK	50 51 105 104 98
TatCy (Bsu) TatCd (Bsu) TatC (Eco) TatCy (Bsu) TatCd (Bsu) TatCd (Eco)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * *  QLPQGSTMIATDVASPFFTPIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR QLTLNAFNLTDPLYVFMQFAFIIGIVLTSPVILYQLWAFVSPGLYEKERKLAVLGPSEILWVYMMLSGICAIAASIPVAAYQLWRFVAPALTKTERK	50 51 105 104 98 155 155
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCy (Bsu) TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * *	50 51 105 104 98 155 155 151
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * * * * * * * * * * *	50 51 105 104 98 155 155 151
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * * *  QLPQGSTMIATDVASPFFTPIKLTFMVSLILSAPVILYQVWAFIAPALYKHERR QLTLNAFNLTDPLYVFMQFAFIIGIVLTSPVILYQLWAFVSPGLYEKERKLAVLGPSEILWVYMMLSGICAIAASIPVAAYQLWRFVAPALTKTERK	50 51 105 104 98 155 155 151
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * * * * * * * * * * *	50 51 105 104 98 155 155 151
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatCd (Bsu)  TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK  * * * * * * * * * * * * * * * * * * *	50 51 105 104 98 155 155 151
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK * * * * * * * * * * * * * * * * * * *	50 51 105 104 98 155 155 151 209 209 205
TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatC (Eco) TatCy (Bsu) TatCd (Bsu)  TatCd (Bsu)  TatCd (Bsu)	MTRMKVNQMSLLEHIAELRKRLLIVALAFVVFFIAGFFLAKPIIVYLQETDEAK MDKKETHLIGHLEELRRRIIVTLAAFFLFLITAFLFVQDIYDWLIRDLDGK  * * * * * * * * * * * * * * * * * * *	50 51 105 104 98 155 155 151 209 209 205

FIG.\_1B

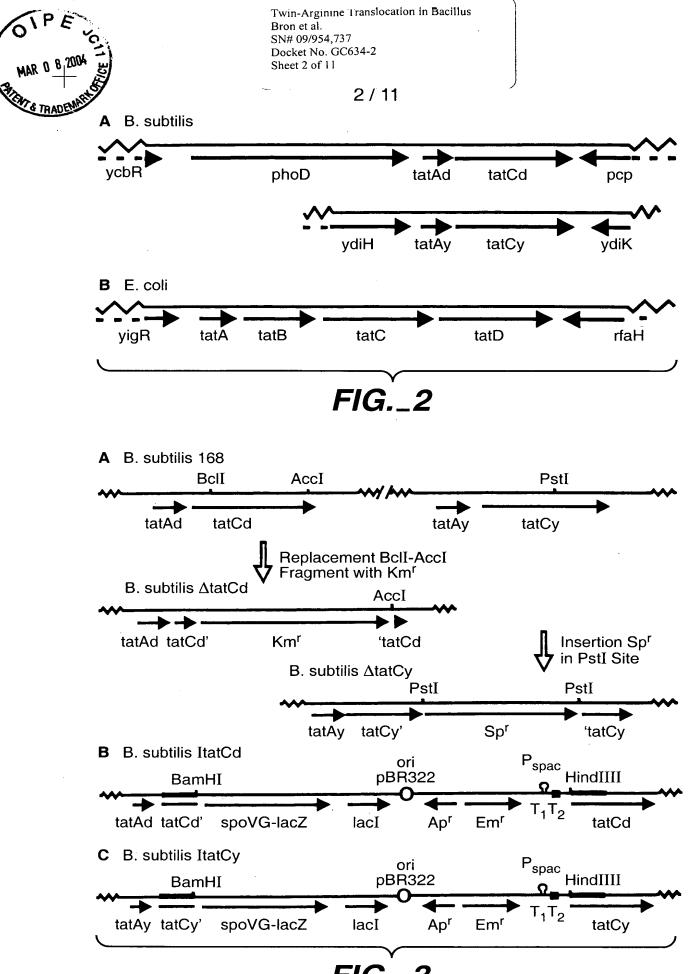


FIG.\_3



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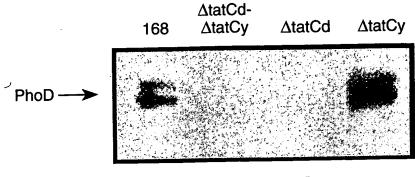


FIG.\_4A

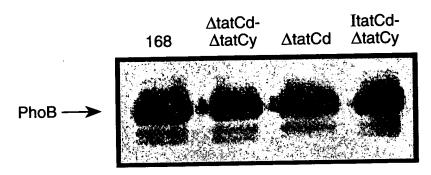
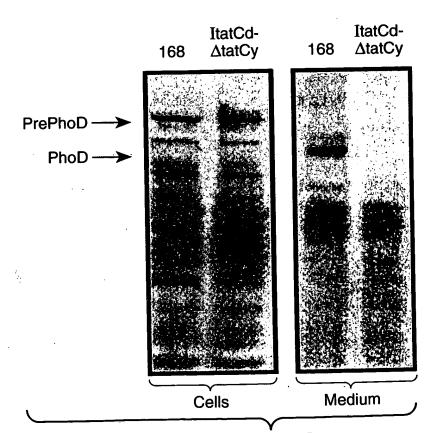


FIG.\_4B





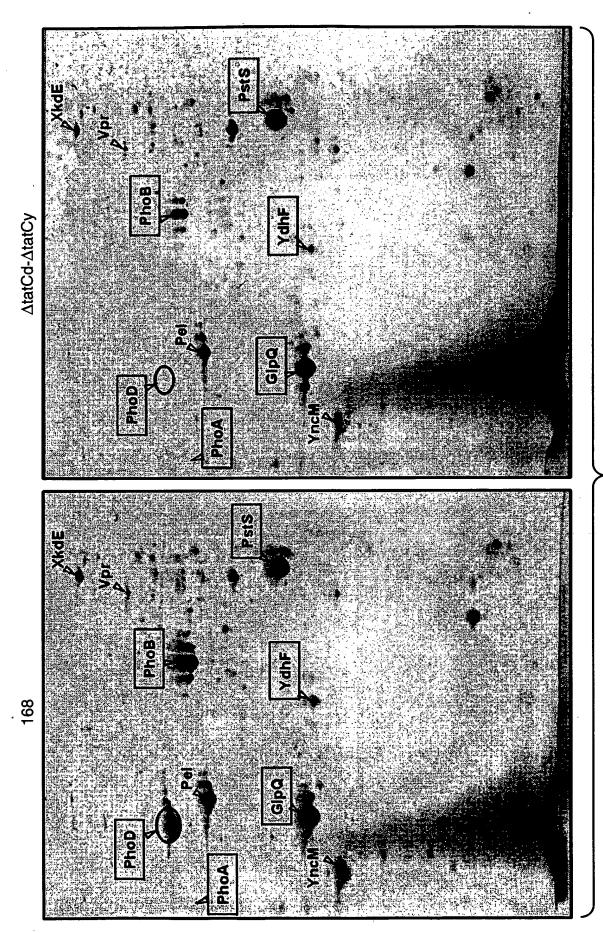
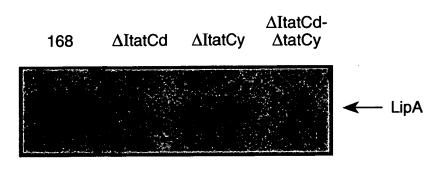


FIG.\_5



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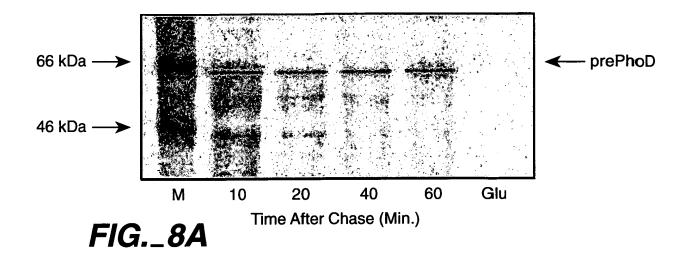


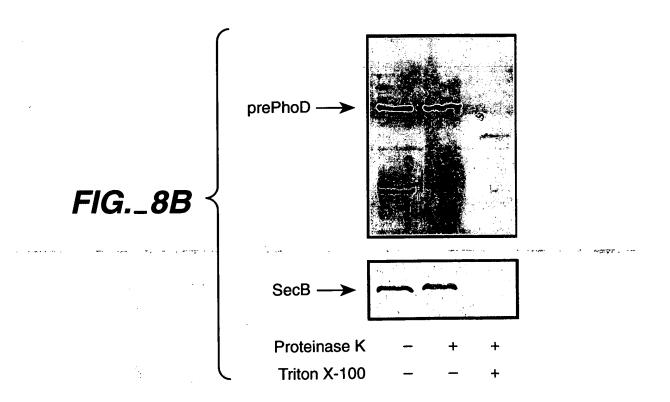
**FIG.\_6** 

Protein	N	h	RR-Motif	H	h	С
AlbB	1	0.1	RRILL	27	2.0	AIA
AmyX TM	9	-0.8	RRSFE	15	1.1	••
AppB TM	8	0.5	RRTLM	19	2.3	-
LipA	7	-1.1	RRIIA	19	1.2	AKA
OppB TM	8	-0.6	<b>RR</b> L <b>V</b> Y	24	2.0	-
PbpX	2	-2.2	<b>RR</b> RK <b>L</b>	14	2.9	WNA
PhoD	3	-1.3	RRKFI	17	0.9	VGA
QcrA TM	1	-1.1	RRQFL	19	1.3	-
TlpA TM	1	-0.8	RRLII	21	2.4	-
WapA W	1	-3.0	RRNFK	18	2.3	VLA
WprA	8	-1.7	<b>RR</b> K <b>FS</b>	20	1.9	AAA
YceA TM	1	-0.4	RRAFL	21	2.2	-
YesM TM	1	-1.5	RRMKI	20	2.4	. QYA
YesW	1	-1.3	RRSCL	19	2.0	VKA
YfkN TM	1	-1.2	RRTHV	17	1.7	IHA
YkpC	8	-1.0	RRVAI	17	2.3	SLA
YkuE	1	-1.3	RRQFL	17	1.0	GYA
YmaC	7	0.0	RRFLL	15	2.4	YSL
YubF ™	9	-2.7	RRNTV	23	2.0	-
YuiC	8	0.2	RRLLM	20	1.9	IEA
YvhJ TM	2	-1.7	RRKIL	18	2.5	-
YwbN	1	-1.8	RRDIL	23	1.4	QTA



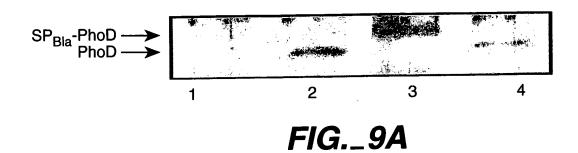
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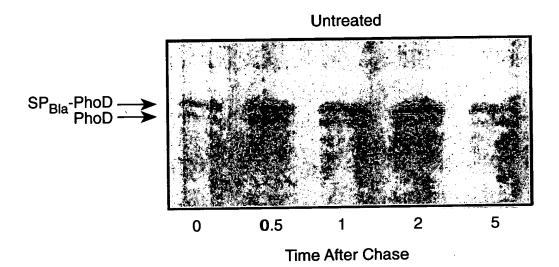


FIG.\_9B

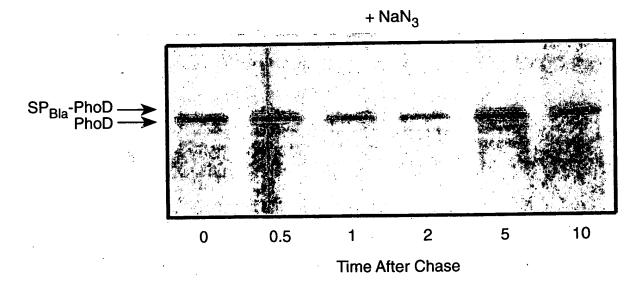
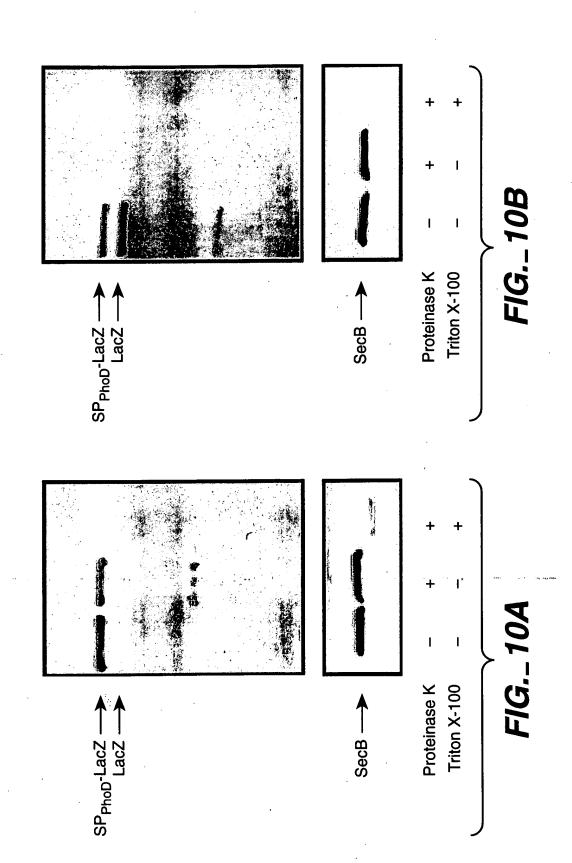


FIG.\_9C

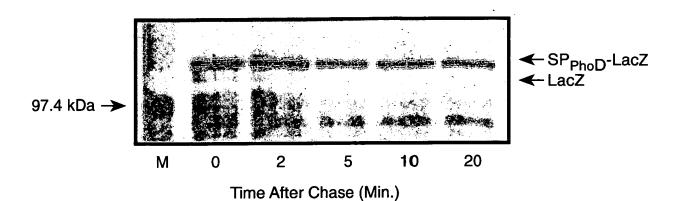


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**FIG.\_11A** 

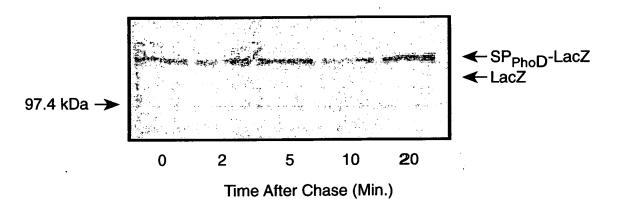
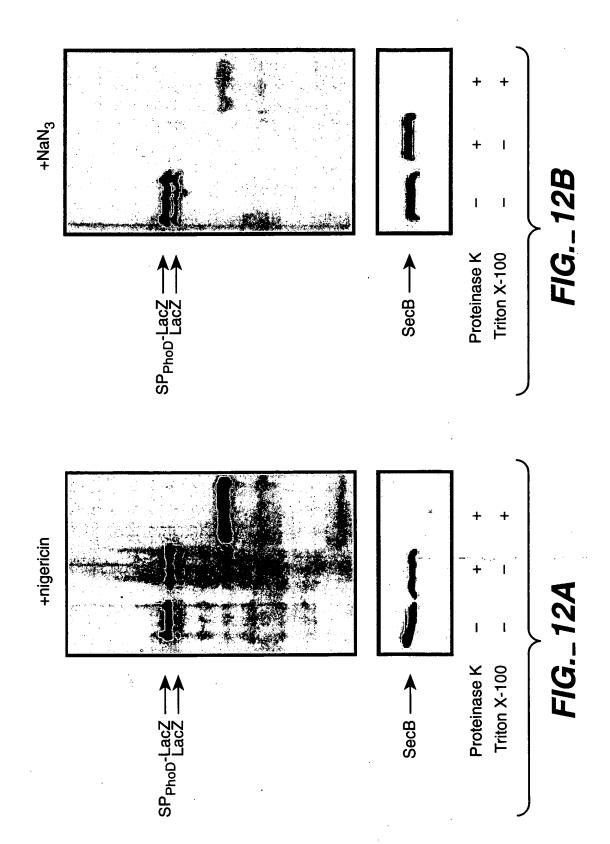


FIG.\_11B



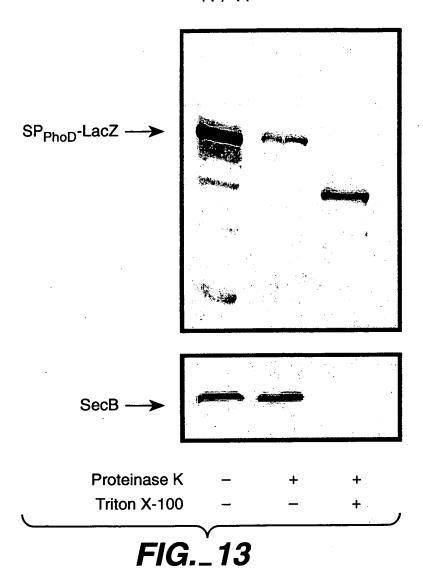
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Homologs in B. alcalophilus

TatA MGGLSVGSVVLIALVALLIFGPKKLPELGKAAGSTLREFKNATK GLADDDDDTKSTNVQKEKA

TatC
MTMMTPNQQTSKKKKRKGRKGRVPMQDMSIMDHAEELRRRIF
VVLAFFIVALIGGFFLAVPVITFLQNSPQAADMPFNAFRLTDPLRV
YMNFAVITALVLIIPVILYQLWAFVSPGLKENEQKATLAYIPIAFL
LFLAGIAFSYFILLPFVISFMGQMADRLEINEMYGINEYFSFLFQL
TIPFGLLFQLPVVVMFLTRLGVVTPTFLRKIRKYAYFALLVIAGII
TPPELTSHLFVTVPMLILYEISITISAITYRKYHGTTDHNGQESAK